

1
SEQUENCE LISTING

<110> SmithKline Beecham Corporation
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Collins, Jon

<120> CAR LIGAND-BINDING DOMAIN POLYPEPTIDE CO-CRYSTALLIZED WITH A
LIGAND, AND METHODS OF DESIGNING LIGANDS THAT MODULATE CAR
ACTIVITY

<130> PR60235

<150> 60/488,415
<151> 2003-07-18

<160> 7

<170> PatentIn version 3.2

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<213> Homo sapiens

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agtctgtgac agccacccca acacgtgacg tc atg gcc agt agg gaa gat gag 293
Met Ala Ser Arg Glu Asp Glu
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ctg agg aac tgt gtg gta tgt ggg gac caa gcc aca ggc tac cac ttt 341
Leu Arg Asn Cys Val Val Cys Gly Asp Gln Ala Thr Gly Tyr His Phe
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aat gcg ctg act tgt gag ggc tgc aag ggt ttc ttc agg aga aca gtc 389
Asn Ala Leu Thr Cys Glu Gly Cys Lys Gly Phe Phe Arg Arg Thr Val
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agc aaa agc att ggt ccc acc tgc ccc ttt gct gga agc tgt gaa gtc 437
Ser Lys Ser Ile Gly Pro Thr Cys Pro Phe Ala Gly Ser Cys Glu Val
40 45 50 55
agc aag act cag agg cgc cac tgc cca gcc tgc agg ttg cag aag tgc 485
Ser Lys Thr Gln Arg Arg His Cys Pro Ala Cys Arg Leu Gln Lys Cys
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Ala	Met	Ala	Leu	Phe	Ser	Pro	Asp	Arg	Pro	Gly	Val	Thr	Gln	Arg	Asp			
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 Gln Glu Ile Cys Ser
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 Phe Ala Gly Ser Cys Glu Val Ser Lys Thr Gln Arg Arg His Cys Pro
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 Ala Cys Arg Leu Gln Lys Cys Leu Asp Ala Gly Met Arg Lys Asp Met
 65 70 75 80
 Ile Leu Ser Ala Glu Ala Leu Ala Leu Arg Arg Ala Lys Gln Ala Gln
 85 90 95
 Arg Arg Ala Gln Gln Thr Pro Val Gln Leu Ser Lys Glu Gln Glu Glu
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 Leu Ile Arg Thr Leu Leu Gly Ala His Thr Arg His Met Gly Thr Met
 115 120 125
 Phe Glu Gln Phe Val Gln Phe Arg Pro Pro Ala His Leu Phe Ile His
 130 135 140
 His Gln Pro Leu Pro Thr Leu Ala Pro Val Leu Pro Leu Val Thr His
 145 150 155 160
 Phe Ala Asp Ile Asn Thr Phe Met Val Leu Gln Val Ile Lys Phe Thr
 165 170 175

4

Lys Asp Leu Pro Val Phe Arg Ser Leu Pro Ile Glu Asp Gln Ile Ser
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Leu Leu Lys Gly Ala Ala Val Glu Ile Cys His Ile Val Leu Asn Thr
 195 200 205

Thr Phe Cys Leu Gln Thr Gln Asn Phe Leu Cys Gly Pro Leu Arg Tyr
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Thr Ile Glu Asp Gly Ala Arg Val Gly Phe Gln Val Glu Phe Leu Glu
 225 230 235 240

Leu Leu Phe His Phe His Gly Thr Leu Arg Lys Leu Gln Leu Gln Glu
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Pro Glu Tyr Val Leu Leu Ala Ala Met Ala Leu Phe Ser Pro Asp Arg
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Pro Gly Val Thr Gln Arg Asp Glu Ile Asp Gln Leu Gln Glu Glu Met
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Ala Leu Thr Leu Gln Ser Tyr Ile Lys Gly Gln Gln Arg Arg Pro Arg
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Asp Arg Phe Leu Tyr Ala Lys Leu Leu Gly Leu Leu Ala Glu Leu Arg
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ctg	gcc	cct	gtg	ctg	cct	ctg	gtc	aca	cac	ttc	gca	gac	atc	aac	act		192
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Tyr	Ile	Lys	Gly	Gln	Gln	Arg	Arg	Pro	Arg	Asp	Arg	Phe	Leu	Tyr	Ala		
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 Arg Ser Leu Pro Ile Glu Asp Gln Ile Ser Leu Leu Lys Gly Ala Ala
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 Gly Thr Leu Arg Lys Leu Gln Leu Gln Glu Pro Glu Tyr Val Leu Leu
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 Tyr Ile Lys Gly Gln Gln Arg Arg Pro Arg Asp Arg Phe Leu Tyr Ala
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